

50 narrow. For example, in MMDA^{-/-} mice release of steel-enclosed Kit-ligand (cKitL) and hematopoietic stem cell motility are impaired resulting in failure of hematopoietic recovery and increased mortality. As described herein, MMDA (matrix metalloproteinase-9/gelatinase B) enables bone marrow repopulating cells to translocate to a permissive vascular niche favoring differentiation and reconstitution of the stem cell and progenitor cell pool. Thus, the invention is directed to a method for ***recruitment*** of adult stem cells in an animal comprising administering to the animal a protease or an activator of a protease wherein the ***recruitment*** translocates an endogenous population of quiescent non-cycling stem cells to a permissive vascular zone in the animal so that the stem cells can proliferate self-renew, differentiate or mobilize to a target site. The activator can be interleukin-1, thrombopoietin, ***G*** - ***C***, GM-CSF, G-CSF, TNF-α or ***fibroblast*** growth factor-4. The protease can be a matrix metalloproteinase, a collagenase, a gelatinase, a stromelysin, a matriplasmin, a metalloelastase, or a membrane-type matrix metalloproteinase. The method may also be used to induce proliferation of adult stem cells in an animal. A therapeutic method for treating a developmental disorder in a mammal using the same is claimed.

US Pat. Appl. Publ., 77 pp.

CODEN: USXXCO

51 T.8 ANSWER 2 OF 11 CAPLUS COPYRIGHT 2008 ACS on STN
TT Device compositions for the ***recruitment*** of cells to blood
AR contacting surfaces in vivo
Methods and compositions for ***recruiting*** cells circulating in the blood stream of a subject to a blood contacting surface and in particular devices and methods for ***recruiting*** endothelial cells to a blood contacting surface of a prosthesis as well as engineering a self-endothelializing graft in vivo by ***recruitment*** of circulating endothelial progenitor cells (EPCs) to form a neo-endothelium on a prosthetic structure are described. Examples are given for ***recruitment*** of endothelial progenitor cells to a blood contacting surface by ligand interaction and by magnetic interaction in vivo.

US Pat. Appl. Publ., 30 pp.

CODEN: USXXCO

52 T.8 ANSWER 3 OF 11 CAPLUS COPYRIGHT 2008 ACS on STN

TT-17
AR A review TT-17 is a potent proinflammatory cytokine produced mainly by activated memory CD4-T cells. The family of TT-17, a new family of cytokines, is composed of six functionally related members, i.e. TT-17 and TT-17R-R in humans and mice. TT-17 exerts its biological activity as a homodimer. In contrast to the selected expression pattern of this gene, the TT-17 receptor is ubiquitously distributed among diverse tissues and cells. TT-17 induces the secretion of TT-6, TT-8, DCER2, MCD-1 and ***G*** - ***C*** by ***fibroblast*** keratinocytes, epithelial and endothelial cells and is also able to induce ICAM-1 expression, T cell proliferation and growth and differentiation of CD34+ human progenitors into neutrophils. The involvement of TT-17 in the rejection of allogeneic grafts has been demonstrated. The potent inflammatory actions that have been identified for TT-17 and the emerging associations with major human diseases such as rheumatoid arthritis and allergic asthma suggest that the family of TT-17 may have significant roles in the pathophysiology of inflammatory processes. TT-17 induces production of metalloproteinases and nitric oxide responsible for the aggravation of arthritis and joint destruction. TT-17 can ***recruit*** and activate neutrophils in the airways mediated by TT-8 and MCD-2. In addition, TT-17 stimulates human bronchial epithelial cells to release the neutrophil-activating factor TT-6.
Biotechnology (Tokyo, Japan) (***2003***), 17(1), 85-97

CODEN: BITPE9; ISSN: 0914-2223

ANSWER 4 OF 11 REQUESTED COPYRIGHT (c) 2002 The Thomson Corporation on STN
Multiple Mechanisms of SOCS3-Mediated Inhibition of ***G***
- ***G*** - Signaling Are Disrupted by C-Terminal Truncation of ***G***
- ***G*** - ***G*** receptor (G-CSFR) signaling regulates survival,
proliferation and differentiation of myeloid progenitor cells into
neutrophils. Different domains and four conserved tyrosine residues in
the receptor C-terminal (Y701 Y729 Y741 Y761) have been linked to
signal transduction pathways that mediate these effects. Suppressor of
cytokine signaling (SOCS) proteins utilize their SH2 domains to inhibit
signaling from both critical tyrosine residues in receptor chains and
intermediate signaling molecules. Members of the SOCS protein family
implicated in negative feedback during hematopoietic growth factor
signaling are GTS SOCS1 SOCS2 and SOCS3. Whether these proteins
downregulate ***G*** - ***G*** signaling and which domains in the
G-CSFR are involved is still unclear. Also it is unknown whether and how
mutations truncating the G-CSFR found in severe congenital neutropenia
(SCN) patients with a predilection to AMI affect SOCS-mediated
downregulation of ***G*** - ***G*** signaling. We first
investigated which SOCS proteins were capable of inhibiting ***G***
- ***G*** - induced STAT3 activity in HEP293 cells. We found that SOCS1
and SOCS2 but not GTS and SOCS3 inhibited G-CSFR activity. Experiments
with tyrosine "null" (m0) and single tyrosine add-back mutants revealed
that while SOCS1-mediated inhibition of ***G*** - ***G***
signaling was entirely independent of C-terminal G-CSFR tyrosines Y729 is
a major ***recruitment*** site for SOCS3-mediated inhibition of
G - ***G*** responses. The fact that the Y729 motif is
universally lost in SCN-derived ***G*** - ***G*** - m0 mutants
indicates that lack of SOCS3 ***recruitment*** may contribute to the
hematopoietic phenotype of DFT.TA715 expressing cells. We therefore
looked in more detail at the role of SOCS3 in DFT.TA715 signaling. SOCS3
is a major transcriptional target of STAT3. Because G-CSFR-DFT.TA715 lacks
two STAT3 activation mechanisms we performed real time quantitative
RT-PCR on bone marrow cells of G-CSFR-DFT.TA715 mice and wild type
littermates. We found that SOCS3 transcript levels are significantly
reduced in DFT.TA715 compared to WT animals. As expected based on the
fact that SCN-derived G-CSFR mutants lack Y729 G-CSFR-DFT.TA715 also
showed reduced responsiveness to SOCS3-mediated inhibition. However
while SOCS3 provided considerable residual inhibition of m0 at higher
SOCS3 concentrations G-CSFR-DFT.TA715 remained completely refractory to
the effects of SOCS3 suggesting a tyrosine independent mechanism of
action involving the receptor C-terminus. Next to the direct inhibitory
effects via SH2 domains SOCS proteins may downregulate signaling via the
SOCS-box which targets internalized receptors for proteosomal degradation.
G-CSFR-DFT.TA715 is severely hampered in ligand-induced endocytosis and may
therefore be refractory to this mechanism. Experiments to test this
possibility using a mutant of SOCS3 lacking the SOCS-box are ongoing. In
conclusion our data show that C-terminal truncations decrease negative
feedback by SOCS3 in three ways: reduced upregulation of SOCS3, lack of
the major SOCS3 ***recruitment*** site Y729 and loss of the C-terminus
with motifs required for internalization and additional tyrosine
independent ***recruitment*** of SOCS3. The lack of appropriate
negative feedback by SOCS3 therefore likely contributed to the
hematopoietic phenotype of truncated G-CSFR mutants found in SCN
Blood (***November 16 2002***) Vol. 100, No. 11, pp. Abstract No.
2850 print

Meeting Info : 44th Annual Meeting of the American Society of Hematology.
Philadelphia, PA, USA. December 06-10, 2002. American Society of
Hematology.

CODEN: BLOOAW. ISSN: 0006-4971.

T.8
TT ANSWERD 5 OF 11 CADTUS CONVENTION 2008 ACS ON STM
Oncostatin M regulates the synthesis and turnover of gp130 leukemia
inhibitory factor receptor .alpha., and oncostatin M receptor .beta. by
distinct mechanisms.
The cytokine receptor subunits gp130 leukemia inhibitory factor receptor
alpha (T.TRP alpha) and oncostatin M receptor beta (OSMR.beta.)
transduce OSM signals that regulate gene expression and cell
proliferation after ligand binding and activation of the Janus protein
tyrosine kinase/STAT and mitogen-activated protein kinase signal
transduction pathways. No feedback processes are ***recruited***.
These processes attenuate receptor action by suppression of cytokine
signaling and by down-regulation of receptor protein expression. This
study demonstrates that in human lung ***fibroblasts*** or epithelial
cells OSM first decreases the level of gp130 T.TRP alpha and OSMR beta.
by ligand-induced receptor desensitization and then increases the level of the
receptors by enhanced synthesis. The transcriptional induction of gp130
gene by OSM involves STAT2. Various cell lines expressing receptor
subunits to the different interleukin-6 class cytokines revealed that only
T.TRP alpha desensitization is promoted by activated ERK and that desensitization of
gp130 OSMR beta and a fraction of T.TRP alpha involves mechanisms that
are seen from signal transduction. These mechanisms include
ligand-mediated dimerization, internalization and endosomal/lysosomal
desensitization. Proteosomal desensitization appears to involve a fraction of receptor
subunit proteins that are ubiquitinated independently of ligand binding.
Journal of Biological Chemistry (***2001***), 276(50), 47038-47045.

CODEN: JBCHA3; ISSN: 0021-9258

T.8
TT ANSWERD 6 OF 11 BTOTS CONVENTION (c) 2008 The Thomson Corporation on STM
Bacterial stimulation of ***G*** - ***CSF*** expression contributes
to PMN persistence at the site of infection.
Bacterial interactions with airway epithelial cells result in the
recruitment and activation of PMNs through ligation of glycolipid
receptors, stimulation of MAP kinases, NF- κ B and the transcription of
proinflammatory cytokines and chemokines. In the studies described, we
demonstrate that *P. aeruginosa* stimulates epithelial ***G*** -
CSF expression which then serves to promote PMN survival.
G - ***CSF*** is produced by macrophages, lymphocytes and
fibroblasts in response to bacterial stimuli although its
function in the airway is less well characterized. To determine if
G - ***CSF*** contributes to airway inflammatory responses, the
expression of ***G*** - ***CSF*** was monitored in several airway
epithelial cell lines following stimulation by *P. aeruginosa*, *S. aureus*
and deleted mutants. There was no constitutive ***G*** - ***CSF***
expression as detected by RT-PCR but 24 hours following exposure to
either organism there was a significant amount of mRNA and a 10-fold
increase in ***G*** - ***CSF*** production as measured by RT-PCR.
The in vitro studies were confirmed by demonstration of ***G*** -
CSF by immunocytochemistry of the bronchial epithelial cells in
sections of mouse lung 24 hours following infection with either *S. aureus*
or *P. aeruginosa*. The physiological effects of ***G*** - ***CSF***
on PMN apoptosis were monitored using a PMN viability assay to compare the
survival of human PMNs incubated in conditioned media harvested from
epithelial cells stimulated with *P. aeruginosa* or from controls. There
was a 75% increase in PMN survival under conditions in which the PMNs were
exposed to media from *P. aeruginosa* stimulated epithelial cells as
compared with controls. This effect was negated by the addition of anti-
G - ***CSF*** but not anti-GM-CSF. The results suggest that
G - ***CSF*** transcription is upregulated in airway epithelial
cells upon stimulation with *P. aeruginosa*, and that the secreted protein
promotes PMN viability.
Abstracts of the General Meeting of the American Society for Microbiology,
(***2001***) Vol. 101, pp. 115. print.

T.8
ANSWER 7 OF 11 MFDLT-TNF ON STN
Bleomycin stimulates lung ***fibroblasts*** to release neutrophil and monocyte chemotactic activity. We determined whether human lung ***fibroblasts*** might release chemotactic activity for neutrophils (NCA) and monocytes (MCA) in response to bleomycin. The human lung ***fibroblasts*** supernatant fluids were evaluated for chemotactic activity by a blind well chamber technique. Human lung ***fibroblasts*** released NCA and MCA in a dose- and time-dependent manner in response to bleomycin. Checkerboard analysis of supernatant fluids revealed that both NCA and MCA were chemotactic. Partial characterization revealed that NCA was partly heat labile trypsin sensitive and predominantly ethyl acetate extractable. In contrast MCA was partly trypsin sensitive and ethyl acetate extractable. The release of chemotactic activity was inhibited by lipoxygenase inhibitors and cycloheximide. Molecular sieve column chromatography revealed that both NCA and MCA had multiple chemotactic peaks. NCA was inhibited by leukotriene R4 receptor antagonist and anti-TL-8 and ***CSF*** -

CSF The MCA was attenuated by leukotriene R4 receptor antagonist and monocyte chemoattractant protein-1 GM-CSF and TGF-beta. The Leukotriene R4 receptor antagonist and those also inhibited the corresponding m w chemotactic activity generated by column chromatography. The concentrations of TL-8 ***CSF*** - ***CSF*** , monocyte chemoattractant protein-1 GM-CSF and TGF-beta in the supernatant fluids significantly increased in response to bleomycin. These data suggest that lung ***fibroblasts*** may modulate inflammatory cell ***recruitment*** into the lung by releasing NCA and MCA in response to bleomycin.

Journal of Immunology (Baltimore, Md. : 1950), *** (1999 May 15) ***

Vol 162 No 10 pp 6200-8
Journal code: 2985117R. ISSN: 0022-1767.

T.9
ANSWER 8 OF 11 CAPTION NUMBER 2002 ACS ON STN
Expression of multiple angiogenic cytokines in cultured normal human prostate epithelial cells: predominance of vascular endothelial growth factor
The cytokines that regulate angiogenesis in normal and malignant prostate tissue are not well studied. Using an RT-PCR-based screen the authors showed that cultured low-passage normal human prostate epithelial cells (PrECs) express a variety of cytokines which have been shown to have angiogenic and/or endothelial cell-activating properties in various systems. These include vascular endothelial growth factor (VEGF), basic fibroblast growth factor (bFGF), transforming growth factor-.alpha. (TGF-alpha), transforming growth factor-beta (TGF-beta), interleukin-8 (IL-8), tumor necrosis factor-alpha (TNF-alpha), granulocyte-macrophage colony-stimulating factor (GM-CSF) and granulocyte colony-stimulating factor (***G** - ***CSF***). Expression of VEGF, bFGF, GM-CSF ***G** - ***CSF*** TGF-alpha and TNF-.alpha. in these cells was confirmed by immunohistochemical culture medium conditioned by normal human PrECs for periods of up to 96 h were found to contain VEGF, GM-CSF ***G** - ***CSF*** IL-8, TGF-beta 1 and TGF-beta 2 but not TNF-alpha or bFGF as detd by ELISA. Of these, VEGF was by far the most prominently expressed angiogenic cytokine (approx. 2500 pg/ml conditioned medium at 96 h vs. 30 to 100 pg/ml conditioned medium for the other cytokines). PrEC-conditioned medium induced an approx. 2-fold stimulation of [³H]-thymidine incorporation in cultured human umbilical cord endothelial cells (HUVECs) deprived of the endothelial growth factors VEGF and bFGF; this stimulation was abolished

by neutralizing antibodies directed against VEGF but not bFGF. TGF- β administration or deprivation of other angiogenic cytokines for which these cells have receptors suggesting that there is not a hierarchy of cytokines controlling its expression; however, retinoic acid a component of DrFC growth medium was found to modestly suppress VEGF at physiol concn (1 ng/ml). These data suggest that normal DrFCs express a variety of angiogenic cytokines most prominently VEGF to ***recruit*** a supporting vasculature even in culture. The authors' data also suggest that the ability of malignant DrFCs to stimulate angiogenesis may be intrinsic and does not need to be acquired during oncogenesis.

International Journal of Cancer (***1999***), 80(6), 868-874

CODEN: IJCNNA; ISSN: 0020-7136

ANSWER 9 OF 11 MRDT-TFC ON STN

MRDT-TCA/T?

Enhanced detection maintenance and differentiation of primitive human hematopoietic cells in cultures containing murine ***fibroblast*** engineered to produce human steel factor, interleukin-3, and granulocyte colony-stimulating factor

To determine whether the sensitivity of the human long-term culture-initiating cell (LTC-TC) assay could be increased we have evaluated a spectrum of different ***fibroblast*** cell lines for their abilities to influence the number of cells detectable as LTC-TC to influence LTC-TC maintenance and/or to influence LTC-TC differentiation into colony-forming cells (CFU) in cocultures containing various sources of LTC-TC. In a series of initial experiments with highly purified subpopulations of CD34+ cells from normal human marrow no significant difference could be found between any of 3 different murine stromal

fibroblast cells in terms of their support of either LTC-TC detection (CFU production) or maintenance (over a 6-week period) and all were equivalent to primary human marrow feeders (HMF). On the other hand, murine M2-10RA ***fibroblast*** engineered to produce high levels of both human granulocyte colony-stimulating factor (***G*** - ***CSF***) and interleukin-3 (TGF- β , 1 ng and 4 ng/ml, respectively) either alone or mixed 1:1 with ST/ST ***fibroblast*** engineered to produce high levels of soluble Steel factor (SF) with or without production of the transmembrane form of SF (60 and 1 ng/ml, respectively) stimulated the production of up to 20-fold more CFU in LTC of cells from normal human marrow. ***G*** - ***CSF*** -mobilized blood or cord blood when compared with parallel cocultures containing HMF. Limiting dilution analysis of the CFU output from all three sources of LTC-TC showed that most of this increase was due to an ability of the engineered feeders to increase the plating efficiency of the LTC-TC assay (approximately 14-fold for marrow LTC-TC and approximately 4-fold for cord blood or mobilized blood LTC-TC). Analysis of the phenotypic of these additionally

recruited LTC-TC from marrow showed they had the same primitive CD34+CD45RA-CD71- phenotype as conventionally defined LTC-TC. The limiting dilution studies also showed that the average number of CFU produced per LTC-TC was additionally and independently increased to yield values of 18 CFU per LTC-TC in marrow, 28 for LTC-TC in cord blood, and 25 for LTC-TC in ***G*** - ***CSF*** -mobilized blood. Repopulating of cells from primary LTC with different feeders into secondary LTC-TC assays containing the best combination of engineered feeders showed that LTC-TCs maintenance could be significantly enhanced (up to 7-fold as compared with primary cocultures containing HMF). However, this enhancement was still not sufficient to amplify the number of LTC-TC present after 6 weeks above the input value. Thus engineering murine ***fibroblast*** to produce sufficient SF ***G*** - ***CSF*** and TGF- β can markedly enhance the detection as well as the maintenance in vitro of a very primitive population of human progenitor cells present in normal adult marrow mobilized blood and cord blood by providing the most sensitive assay conditions thus far described. The present findings also provide

new evidence of biological heterogeneity between different cell populations that can be operationally identified as T_H-1 or thus re-emphasizing the importance of limiting dilution analyses to distinguish between quantitative and qualitative effects on these cells.

Blood *** (1996 Nov 15) ** Vol 88 No. 10, pp. 3765-73.

Journal code: 7603509. ISSN: 0006-4971.

ANSWER 10 OF 11 MRTD-TNF ON STN

Microenvironmental influences on inflammatory cell differentiation. Airways inflammation involves accumulation of inflammatory cells such as eosinophils, basophils and mast cells which are derived from progenitors in marrow and blood. The inflamed tissue of the airways through its structural (epithelium, stroma) and inflammatory cell components, produces an array of cytokines which can influence the differentiation of, inflammatory cell progenitors. It is particular mechanism that we have investigated showing that molecules such as GM-CSF ***G***

CSF TNF- α and SCF can be produced by airways epithelial cells and ***fibroblasts*** in quantities sufficient to induce hemopoietic events either systemically or locally. Corticosteroids may act therapeutically at least in part to block inflammatory cell differentiation and thus ***recruitment***, into the allergic inflammatory process in the airways.

Allergy *** (1995) ** Vol 50 No 25 Suppl, pp. 25-8. Ref: 20

Journal code: 7804028. ISSN: 0105-4538.

ANSWER 11 OF 11 BIOSIS COPYRIGHT (c) 2008 The Thomson Corporation on STN

CYTOKINES AND FUNGAL INFECTIONS

Cytokines play a major role within the immune system as necessary mediators involved in the set up of the specific immune response as well as factors which amplify the anti-microbial response. It includes activation of B and T lymphocytes leading to the production of antibodies and generation of specific cytotoxic T cells, activation of phagocytic cells, ***recruitment*** of cells at the infectious site and stimulation of the hematopoiesis leading to an increased number of active cells. Furthermore, cytokines are a link between immune cells and many others of the organism. For example some of them act on the central nervous system and possess pro-inflammatory activities. During fungal infections, monocytes and macrophages are triggered either by fungi or their derived products and release tumor necrosis factor alpha (TNF-alpha) and interleukin-1 (IL-1). Other cells such as

fibroblasts NK cells, and neutrophils can be activated by fungi to produce cytokines. In addition, T lymphocytes are also stimulated and release interferon gamma (IFN-gamma), a potent macrophage activator. If an excess of cytokines is produced, it is then possible to detect them in plasma. This has been reported for TNF-alpha, IL-2 and IL-6. The capacity of these cytokines to enhance in vitro anti-fungal activities of phagocytic cells has been reported for TNF-gamma, IL-1 alpha and macrophage-colony stimulating factor (M-CSF) which amplifies the monocytes/macrophages killing activities and for TNF-gamma and IL-8 which trigger the functions of neutrophils. In the mouse model *in vivo*, experiments have demonstrated the efficiency of IL-1, M-CSF, ***G***

CSF and TNF-alpha to increase the protection against fungal infections. In addition, some cytokines like IL-1 can act synergistically with anti-fungal drugs. The efficiency of these cytokines are often associated with a pre-treatment before injection of the microbial agent. Thus, these animal models are not always reliable for the treatment of human fungal infections. Th1 as well as Th2 T lymphocyte subpopulations which differ in their pattern of cytokine production can be activated during experimental fungal infection and lead to different immunity associated or not with protection. Further knowledge of the immune response to fungal infections as well as a better

understanding of the relative role of individual cytokines to induce protection will help in the near future to enhance the defense mechanisms involved during fungal infections.
Journal de Mycologie Medicale, (***1992***) Vol. 2, No. 2, pp. 61-67.
ISSN: 1156-5233.

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T.4 29452 G-CSE
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